Convertable (c) 1993 - 2000 Computer Ltd
OM protein - protein search, using sw model
Run on: December 27, 2001, 16:48:55; Search time 25.08 Seconds (without allgoments) 604.754 Million cell updates/sec
Title: US-09-830-647-1 Perfect score: 3510 Sequence: 1 MNSGAMRIHSKGHFQGGIQVSDNLLTAFFSSPSTSTFTGF 674
Scoring table: BLOSUM62 Gapop 10.0 , Gapext 0.5
Searched: 212252 segs, 22503292 residues
Total number of hits satisfying chosen parameters: 212252
Minimum DB seq length: 0 Maximum DB seq length: 2000000000
Post-processing: Minimum Match 10% Maximum Match 100% Listing first 45 summaries
Database: Issued_Patents_AA:* 1: /cgn2_6/ptodata/2/iaa/5A_COMB.pep:* 2: /cgn2_6/ptodata/2/iaa/5B_COMB.pep:* 3: /cgn2_6/ptodata/2/iaa/5B_COMB.pep:* 4: /cgn2_6/ptodata/2/iaa/6B_COMB.pep:* 5: /cgn2_6/ptodata/2/iaa/PCTUS_COMB.pep:* 6: /cgn2_6/ptodata/2/iaa/PCTUS_COMB.pep:*
Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
SUMMARIES Result Query No. Score Match Length DB ID Description
158 4.5 783 6 5231168-2 Patent No. 523116 156.5 4.5 3248 1 US-08-353-700-1 Sequence 1, Appl
130.5 4.5 3248 5 FCT-US95-16216-1 Sequence 1. 55.5 4.4 248.2 1 US-08-328-524-6 Sequence 6. 146 4.2 2391 2 US-08-446-855A-2 Sequence 2.
146 4.2 2391 4 US-09-150-741-2 Sequence 2, Appl 144 4.1 688 3 US-09-140-047-8 Sequence 8, Appl 140 5 4.0 000 1 US-08-425-061-20 Sequence 8, Appl 140 5 4.0 000 1 US-08-425-061-20 Sequence 8, Appl 140 5 4.0 000 1 US-08-425-061-20
140.5 4.0 900 1 US-08-425-061-20 Sequence 20, 140.5 4.0 900 2 US-08-825-886-20 Sequence 20, 140.5 4.0 914 1 US-08-435-661-21 Sequence 21, 140.5 4.0 914 1 US-08-435-661-21
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140.5 4.0 1363 1 US-08-425-061-23 Sequence 23, 140.5 4.0 1363 2 US-08-825-886-23 Sequence 23,
140.5 4.0 1852 1 US-08-425-061-24 Sequence 24, 140.5 4.0 1852 2 US-08-825-886-24 Sequence 24,
140.5 4.0 1863 1 US-08-425-061-16 Sequence 2
140.5 4.0 1863 1 US-08-483-553-2 Sequence 2,
140.5 4.0 1863 1 US-08-487-002-2 Sequence 2, 140.5 4.0 1863 1 US-08-483-554B-2 Sequence 2,
140.5 4.0 1863 1 US-08-488-011B-2 Sequence 140.5 4.0 1863 2 US-08-825-886-16 Sequence 140.5 4.0 1863 4 US-08-850-727-2 Sequence
140.5 4.0 1863 5 PCT-US95-10202-2 Sequence 2, 140.5 4.0 1863 5 PCT-US95-10203-2 Sequence 2,

RESULI 52311 ; Pate ; Pate; VUUUS; ; VUUUS; ; SEQ ; SEQ Qy Qy Db	
RESULT 1 5231168-2 ; Patent NG	4444 544 544 544 544 544 544 544 544 54
S. 52: S. 62: S. 62: S. OF S. S. N. S.	140.5 140 140 140 140 140 140 140 140 140 140
5231168 ANT: DZIEGIEL, MOR S; RIENECK, KLAUS; W OF INVENTION: MALA OF INVENTION MALA OF PEDICATION DATA ICATION NUMBER: U MG DATE: 18-SEP-19 2: 15; Conservative LGGRVEEFLSKDISYLIS LGGRVEEFLSKDISYLIS LGGRVEEIVEQUNDES SWGVKILHIDDI-RYYIE EGG-FEBAHHENFSSEVS FKSPDTVCLSR FKSPDTVCLSR LIGHT: :: : : : ESWGVKILHIDDI-RYYIE SWGVKILHIDDI-RYYIE LIGHT: :: : : : ENFEKIEFSUQLKEK KPFVKVEDMSQLYRPFYL LILL: : : : : KPFEEIEKVDVQPKIVDL KWGTTSIQLQLKEK LILL: : : : : : EXPFEEIEKVDVQPKIVDL KYGGTSIQLQLKEK LILL: : : : : : EDDKNEKVEHEIVEVEEI EDDKNEKVEHEIVEVEEI EDDKNEKVEHEIVEVEEI ETEKKILFISEPIPHPSN : : : : : : : : : : : : : : : : : : :	444446600000000000000000000000000000000
31168 31168 31168 31168 31168 31168 31168 31168 31168 31168 31170N: MALARIA ANTIGE SEQUENCE: 19 SEQUENCE: 19 PELICATION DATA: TION NUMBER: US/07/409,6 DATE: 18-SEP-1989 783 783 783 783 783 783 783	1863 1863 1863 1863 1863 1863 1863 1863
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. 5231168 CANT: DZIEGIEL, MORTEN; BORRE, MARTIN; JEPSEN, SOREN; NS; RIEMECK, KLAUS; WIND, ANNETTE; JAKOBSEN, PALLE H. OF INVENTION: MALARIA ANTIGEN OF INVENTION: MALARIA ANTIGEN R OF SEOUBLCES: 19 ROF SEOUBLCES: 19 ROF SEOUBLCES: 19 ROF SEOUBLES: 105/07/409,658 ILCATION NUMBER: US/07/409,658 IXIGATION UNMBER: US/07/409,658 IXIGATION AUMBER: US/07/4	PCT-US95-10220-2 US-08-598-591-2 US-08-798-691-2 US-08-798-691-4 US-08-825-487A-2 US-08-825-487A-2 US-08-825-487A-6-2 US-09-074-476-4 US-09-074-476-6 US-09-074-476-6 US-09-074-476-6 US-09-08-487-826B-4 US-08-588-459A-4 US-08-687-080-552-126-14 US-08-687-080-552-126-14 US-08-687-080-558-588-592-126-14 US-08-687-080-558-588-592-126-14 US-08-687-080-552-126-14 US-08-687-080-552-126-14 US-08-687-080-552-126-14
JAKOBSEN, PALLE H. JAKOBSEN, PALLE H. JAKOBSEN, PALLE H. 6.4e-05; 6.4e-05; SGRISPVPSPESAYTAETTSP: : : : : DKSVTE-PAEHEEVVSEESNP IDF	11-2 11-2 11-2 11-4 11-6 11-6 11-6 11-6 11-6 11-6 11-6
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4; Gaps	0004404404004000
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; MOLECULE TYPE: P
; HYPOTHETICAL: NO
; ANTI-SENSE: NO
; ORIGINAL SOURCE:
; ORGANISM: HUMA
US-08-353-700-1
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US-08-353-700-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (215) 563-4044
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENCTH: 3248 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: REED, JANET E.
REGISTRATION NUMBER: 36,252
REGISTRATION NUMBER: 4100:
TELEPHONE: (215) 563-4044
TELEPHAX: (215) 563-4044
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
2505 SEL-----EIINSSFENILQEKEQEK------VQMKEKSSTAMEMLQTQLKELNER 2549
                                                                                                                                                                             2457 NSKGEVE-TLKAKIEGMTQSLRGLELDVVTIRSEKENLTNEL------QKEQERI 2504
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
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APPLICANT: RATTNER, JEROME B.

TITLE OF INVENTION: NUCLEIC ACID ENCODING A

TITLE OF INVENTION: TRANSIENTLY EXPRESSED KINETOCHORE PROTEIN,

TITLE OF INVENTION: AND METHODS OF USE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  495 GOHEIVEV-EEILPEDKNE--KVOH-----EIVEVEEILPEDKNEKGOHEIVEVEEILP 545
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                                                                                 69 KDLGGRVEEFLSKDISYLISNKKEAKFAQTLGRISPVPSPESAYTA-----E 115
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TOPOLOGY: 11
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                                                                                                                                                                                                                                                                                                                                                                                                  Local
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RESULT 3
PCT-US95-16216-1
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/353,700
FILING DATE: 09-DEC-1995
ATTORNEY/AGENT INFORMATION:
NAME: Reed, Janet E.
REGISTRATION NUMBER: 36,252
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DDS/MS-DOS
SOFTWARE: PAt-entin Release #1.0/
CURRENT APPLICATION DATA:
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APPLICANT: Rattner, Jerome B.

TITLE OF INVENTION: Nucleic Acid Encoding a Transiently
TITLE OF INVENTION: Expressed Kinetochore Protein, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Dann, Dorfman, Herrell and Skillman
STREET: 1601 Market Street Suite 720
CITY: Philadelphia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2959 KYCSLLISHEKL-----
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2697 KMDKMSFVE-KVNKMTAKETELQREMHEMAQKTAELQ------EELSGEK 2739
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2648 KEQNLELRNLTVELEQKIQVLQS-----KNASLQDTLEV---LQS---SYKNLENELELT 2696
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2604 QEVEDGKQKLEKKDEEISRLK---NQIQDQEQLVSKLSQ-------VEGEHQLW 2647
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2550 VAALHNDQEACKAKEQNLSSQVECLE-----LEKAQLLQGLDEAKNNYIVLQSSVKGL-I 2603
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                398 LYK-ETQETEKK-LLFISEPIPHPSNELRGLNEKMSNKCSMLSTAEDDIRQNFTQLPLHK 455
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       349 KR-----IKYSVGSLSPVSASVLKKTEQKEKVELQHISQKDCQEDDTTVKEQNF 397
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                                                                                                                                                                                           CLASSIFICATION:
                                                                                                                                                                                                                       APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EKKKKGYCECCLQKYEDLETHLLSEQHRNFAQSNQYQVVDDIVSKLVFDFVEYEKDTPKK 348
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                                                                                                                                                                                                                                                                                                      Release #1.0, Version #1.30
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US-08-328-254-6
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PCT-US95-16216-1
Sequence 6, Application US/08328254
Patent No. 5710022
GENERAL INFORMATION:
APPLICANT: Zhu, Xueliang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE: (215) 563-4100
TELEFAX: (215) 563-4044
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                   2994 RGSPLLGPVVPGPSPIPSVTEK 3015
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                                                                                                                                                                                                           EKSEFLG-----FTSYTEK 640
                                                                                                                                                                                                                                                                                                                                       LEEEKEILQKELSQLQAAQEKQKTGTV-MDTKVDELTTEIKELKETLEEKTKEAD-EYLD 2958
                                                                                                                                                                                                                                                                                                                                                                                                                         SKEELNNSLKATTQILEELKKTKMDNLK-----YVNQLKKENERAQGKMKLLIKSCKQ 2900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NKQECILDISEHTLSENDLEELRVDHYKCNIQASVHVSDFSTDNSGSQPKQ----KSDTV 511
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EYQLRLHEAEKKHQALLLDTNKQYEVEIQTYREKLTSKEECLSSQKLEIDL-----LKS
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                                                                                                                                                                                                                                                       KYCSLLISHEKL-----
                                                                                                                                                                                                                                                                                           SLPSGKI-HRKVKIILGRNRKENLEPNAEFDKRTEFITQEENRICSSPVQSLLDLFQTSE 626
                                                                                                                                                                                                                                                                                                                                                                            LFPAKDLKEKDLHSIFT----HDSGLITINSSQEHLTVQAKAPFHTPPEEPNECDFKNMD 567
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LYK-ETQETEKK-LLFISEPIPHPSNELRGLNEKMSNKCSMLSTAEDDIRQNFTQLPLHK 455
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NRLAGELQLLLEEIKSSKDQLKELT---LENSELKKSLDCMH----KDQVEKEGKVREEIA 2793
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (619) 535-894
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: P-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 535-9001
TELEFAX: (619) 535-8949
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 22-OCT-1993
ATTORNEY/AGENT INFORMATION:
NAME: Campbell, Cathryn
31,
                                                                         2012
                                                                                                                                                     1969
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: A NUMBER OF SEQUENCES: 8 CORRESPONDENCE ADDRESS:
2066 EYQLRLHEAEKKHQALLLDTNKQYEVEIQTYREKLTSKEECLSSQKLEIDL---
                                                                                                                                                                                                                                                                                                       1876 QEVEDGKQKLEKKDEEISRLK---NQIQDQEQLVSKLSQ
                                                                                                                                                                                                                                                                                                                                                                                 1822 VAALHNDQEACKAKEQNLSSQVECLE-----LEKAQLLQGLDEAKNNYIVLQSSVNGL-I 1875
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1729 NSKGEVE-TLKAKIEGMTQSLRGLELDVVTIRSEKENLTNEL------QKEQERI 1776
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                                                                                                                                                                                                                                                                                                                                             170
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                                                                                                                                                                                                                                                                                                                                                                                                                     116 TTSPHPSHDGSSFK----SPDTVCLSRGKLLVEKA--IKDHDFIPSNSILSNALSWGVKI 169
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/08/328,254 FILING DATE: 24-OCT-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 69 KDLGGRVEEFLSKDISYLISNKKEAKFAQTLGRISPVPSPESAYTA-----E 115
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REGISTRATION NUMBER:
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                                 LYK-ETQETEKK-LLFISEPIPHPSNELRGLNEKMSNKCSMLSTAEDDIRQNFTQLPLHK 455
                                                                                                                                                                                                                             KEQNLELRNLTVELEQKIQVLQS-----KNASLQDTLEV---LQS---SYKNLENELELT 1968
                                                                                                                                                                                                                                                                                                                                           LHIDDIRYYIEQKKKELYLLKKSSTSVRDGGKRVGSGAQKTRTGRLKKPFVKVEDMSQLY 229
                                                                                                                                                     KMDKMSFVE-KVNKMTAKETELQREMHEMAQKTAELQ-----EELSGEK
                                                                                                                                                                                 EKKKKGYCECCLOKYEDLETHLLSEQHRNFAQSNQYQVVDDIVSKLVFDFVEYEKDTPKK 348
                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEL----EIINSSFENILQEKEQEK------VQMKEKSSTAMEMLQTQLKELNER 1821
                                                                                                                                                                                                                                                               RPFYLQLTNMPF-INYSIQKPCSPFDVDKPSSMQKQTQVKLRIQTDGDKYGGTSIQLQLK 288
                                                                         NRLAGELQLLLEEIKSSKDQLKELT---LENSELKKSLDCMH----KDQVEKEGKVREEIA 2065
                                                                                                           KR------IKYSVGSLSPVSASVLKKTEQKEKVELQHISQKDCQEDDTTVKEQNF 397
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: California
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4370 La Jolla Village Drive, Suite 700
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; TELEPHONE: 703-816-4000
; INFORMATION FOR SEQ ID NO: 2:
; SEQUENCE CHARACTERISTICS:
LENGTH: 2391 mnino acids
; TYPE: amino acid
; TYPE: amino acid
; STRANDEDNESS: single
; MOLECULE TYPE: protein
US-08-446-855A-2
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Sequence 2, Application US/08446855A
Patent No. 5849573
GENERAL INFORMATION:
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Query Match
17.6%; Prod. No. 0.0034;
Matches 129; Conservative 139; Mismatches 264; Indels 202; Gaps
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APPLICATION NUMBER: US/08/446.
FILING DATE: 06-7ul-1-995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Mitchard, Leonard C
REGISTRATION NUMBER: 29.09
REGISTRATION NUMBER: 47-8
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-816-4000
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APPLICANT: Flores, Maria V
APPLICANT: o'Sullivan, William J
TITLE OF INVENTION: Nucleotide sequence encoding carbamoyl
TITLE OF INVENTION: phosphate synthetase II
NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-POS
SOFTWARE: PATENTIN Release #1.24
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1177 DKQIAHYLSFNTSDNNNNNNNISSCRVTENDVMKYREKLGLFPHIKVIDTLSAEFPALTN 1236
                                                                                                1119 QAFHLNMPMDKIHELTHIDYWFLHKFYNIYNLQNKLKTL--KLEQLSFNDLKYFKKHGFS 1176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION DATA:
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                                              89 NKKEA-----KFAQTLG---RISPVPSPESAYTAETT 117
                                                                                                                                                  49 KVFYLDLPSVTISE------KLQKDIKDLGGRVEEFLSKDISYL----IS 88
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1100 No. 5849573th Glebe Road, 8th Floor
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CURRENT APPLICATION NUMBER: US/09/150,741
CURRENT FILING DATE: 1998-09-10
EARLIER APPLICATION NUMBER: PL6380
EARLIER FILING DATE: 1992-12-16
EARLIER FILING DATE: 1993-12-02
EARLIER APPLICATION NUMBER: AU93/00617
EARLIER FILING DATE: 1993-12-02
EARLIER APPLICATION NUMBER: 08/446.855
EARLIER FILING DATE: 1995-07-06
NUMBER OF SEQ ID NOS: 15
SOFTWARE: PATENT NOS: 15
SEQ ID NO 2
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TITLE OF INVENTION: Nucleotide Sequence Encoding Carbamoyl Phosphate
: Patent No. 6183996
; TITLE OF INVENTION: Synthetase II
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US-09-150-741-2
; Sequence 2, Application US/09150741
; Patent No. 6183996
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; OTHER INFORMATION: Description of Artificial Sequence: protein 05\text{-}09\text{-}150\text{-}741\text{-}2
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                                                          TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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                                                                                             GENERAL INFORMATION:
APPLICANT: Walker, David H.
APPLICANT: YU, XUE-Jie
TITLE OF INVENTION: Ehrlichia canis
TITLE OF INVENTION: Protein Gene
                                                                                                                                                                                                       Sequence 8, Application US/09141047A Patent No. 6043085
FILE REFERENCE: D6143
CURRENT APPLICATION NUMBER: US/09/141,047A
CURRENT FILING DATE: 1998-08-27
NUMBER OF SEQ ID NOS: 15
SEQ ID NO 8
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          118 SPHPSHDGSSFKSPDTVCLSRGKLLVEKAIKDHDFIPSN-----SILSNALSWGVKILH 171
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       49 KVFYLDLPSVTISE------KLOKDIKDLGGRVEEFLSKDISYL----IS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity les 129; Conserv
                                                                                                                                                                                                                                                                                                                                                                      NLLTAFFSSPSTST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EYNNNV - - NMNCSNYNNASAFVNGKDRNDNL - - - ENDCIEKNMDHTYKHYNRLNNRRSTN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HYKCNIQASVHVSDFSTDNSGSQPKQKSDTVLFPAKDLKEKDLHSIFTHDSGLITINSSQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NTSDEEIGNN-KNMDMYLSKEKSISNKNPGNSYYVVDSVYNNEYKINKMKELIDNENLND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NIN----DECKTYKK------NKYKDMGLNNNTVDELSNGTSHSTNDHLYLDNF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EQNFLYKETQETEKKLLFISEPIPHPSNELR--GLN----EKMSNKCS------ML
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NTTTYFMEGQNSVINNKNKENSSLLKGDEEDIVMVNLKKENNYNSVINNVDCRKKDMDGK 1492
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              KYS------VGSLSPVSASVLKKTEQ-----KEKVELQHISQKDCQEDDTTVK 393
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SIG-----SDDTNILSAQNSN----NNFSCNNENMNKANVDVNVLENDIKKREDI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            KKGYCECCLQKYEDLETHLLSEQHRNFAQSNQYQVVDDIVSKLVFDFVEYEKDTPKKKRI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NNINIENNICHDISINKNIKVTINNSNNSISNNENVETNLNCVSERAG--SHHIYGKEEK 1386
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FYLQLTNMPFINYSIQKPCSPFDVDKPSSMQKQTQVKLRIQTDGDKYGGTSIQLQLKEKK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    YLYLTYQG------QEHDVLPLNMKRKKICTLNNKRNANKKKYH
                                                                                                                                                                                                                                                                                                                                                                                                                                                 KRTEFIT-QEENRICSSPVQSLLDLFQTSEEKSEFLGFTSYTEKSGICNVLDIWEEENSD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ERMMLMV - - - - NNEKESNHE-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EHLTVQAKAPFHTPPEEPNECDFKNMDSLPSGKIHRKVKIILGRNRKENLEPNA---EFD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STAEDDIRQNFTQLPLHKNKQECILD------ISEHTLSENDLEEL-----RVD
                                                                                                                                                                                                                                                                                                                                                                                                               KNYHYVNHKRNNEYNSNNIESKFNNYVDDINKKEY----YEDENDIYYFTHSSQGNNDD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           IDDIRYYIEQKKKELYLLKKSSTSVRDGGKRVGSGAQKTRTGRLKKPFVKVEDMSQLYRP 231
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative 139;
                                                                                                                                                                                                                                                                                                                                                                      670
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4.2%; Score 146; DB 4; Length 2391; 17.6%; Pred. No. 0.0034;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           -----KFAQTLG----RISPVPSPESAYTAETT
                                                                                                                         120-kDa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           KGHRR-NGLNKKNKEKNMEKNKGKNKDK 1693
                                                                                                                         Immunodominant Antigenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      264;
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                                                                                                                                                                                                                                                                                                                                                                                                               1748
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US-08-425-061-20 US-08-425-061-20 ; Sequence 20, Application ; Patent No. 5622829 ; GENERAL IMFORMATION:

Application US/08425061

APPLICANT:
APPLICANT:
APPLICANT:

ROWELL, Sarah LYNCH, Eric SZABO, Csilla APPLICANT:
APPLICANT:

KING, Mary-Claire FRIEDMAN, Lori OSTERMEYER, Beth

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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Ehrlichia canis
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT
 571
                                   637
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    53 LDLPSVTISEKLQKD------IKDLGGRVEEFLS-KDISYLISNKKEAKFAQTLGRI 102
KEEST 575
                              YTEKS 641
                                                                                                                                                                    TVQAKAPFHTPPEEPNECDFKNMDSLP-----SGKIHRKVKIILGRN-----RKEN
                                                                                                                                                                                                                                   CNIQASVHVSDFSTDNSGSQPKQKSDTVLFPAKDLKEKDLHSIFTHDSGLITINSSQEHL
                                                                                                                                                                                                                                                                                                     NEKMSNKCSMLSTAE---DDIRONFTQLPLHKNKQECILDISEHTLSENDLEELRVDHYK 483
                                                                                                                                                                                                                                                                                                                                                                                                                           VDGSIEHSSSEVGEKVSKTSKEESTPEVKAEDLQPAVDDSVEHSSSEV----GEKVSETSK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            --NSILSNALSWGVKILHIDDIRYYIEQKKKELY-----LLKKSSTSVRDGGKRVGSGAQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SPVPSPESAYTAETTSPHPSHDGSSFKSPDTVCLSRGKLLVEKA-----IKDHDFIPS 155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  IDNNNVTTSSTQDKSGNLMEVIMRILNFGNNSDEKVSNEDTKVLVESLQPAVNDNVGNPS
                                                                 LQPAVDGSVEHSSSEVGEKVSETSKEESTPEVKAEDLQPAVDGSVEHSSSEVGEKVSETS
                                                                                                                                  SSEVGEKVSETSKEENTPEVKAEDLQPAVDGSVEHSSSEVGEKVSETSKEESTPEVKAED
                                                                                                                                                                                                     PAVDGSVEHSSSEVGEKVSETSKEENTPEVKAEDLQP------AVDGSVEHS
                                                                                                                                                                                                                                                                     GEKVSETSKEESTPEVKAEDL-QPAVDSSIEHSSSEVGKKVSE-TSKEESTPEVKAEDLQ 404
                                                                                                                                                                                                                                                                                                                                        LQPAVDDSVEHSSSEVGEKVSETSKEEN----TPEVRAEDLQPAVDGSVEHSSSE---V
                                                                                                                                                                                                                                                                                                                                                                        -----LQHISQKDCQEDDTTVKEQNFLYKETQETEKKLL--FISEPIPHPSNELRGL 426
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ----GEKVSKTSKEESTPEVK----AEDLQPAVDDSVEHSSSEVGEKVSETSK----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LRIQTDGDKYGGTSIQLQLKEKKKKGYCECCLQKYEDLETHLLSEQHRNFAQSNQYQVVD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              KTRTGRLKKPFVKVEDMSQLYRPFYLQLTNMPFINYSIQKPCSPFDVDKPSSMQKQTQVK 268
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEVGKEENAPEVKAEDLQPAVDGSVEHSSSEV----GKKVSETSKEESTPEVKAEDLQPA 118
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       133;
                                                                                                 LEP-----NAEFDKRTEFITQEEN--RICSSPVQSLLD--LFQTSEEKSEFLGFTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4.1%; Score 144; DB 3; ilarity 20.0%; Pred. No. 0.0008; Conservative 104; Mismatches 260
                                                                                                                                                                                                                                                                                                                                                                                                           ----EENTPEVKAEDLQPAVDGSVEHSSSEVGKKVSETSKEESTPEVKAED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          kDa
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                                                                                                    636
                                                                                                                                                                      589
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; SEQUENCE CHARACTERISTICS:

; LENGTH: 900 amino acids

; TYPE: amino acid

STRANDEDNESS: single

; TOPOLOGY: linear

; MOLECULE TYPE: protein

US-08-425-061-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: OSMAN, Richard A
REGISTRATION NUMBER: 36,627
REFERENCE/DOCKET NUMBER: A-59
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 494-8700
TELEPHONE: (415) 494-8771
TELEPHONE: (415) 494-8771
TELEPHONE: 910 277299
INFORMATION FOR SEQ ID NO: 20:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTMARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/425,061
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: LEE, Ming
TITLE OF INVENTION: GENETIC
TITLE OF INVENTION: CANCER
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
        672 KEPATGAKKSN---KPNEQTSKR------HDSDTFPELKLTNAPGSFTKCSNTSELK 719
                                                     386 QEDDTTVKEQNFLYKETQETEKKLLFISEPIPHPSN---ELRGLNEKMS-NKCSMLSTAE 441
                                                                                                        612 RRKSSTRHIHALELVVSRNLSPPNCTELQIDSCSSSEEIKKKKYNQMPVRHSRNLQLMEG 671
                                                                                                                                                        352 KYSVG------SLSPVSASVL------KKTEQKEKVELQHISQKDC 385
                                                                                                                                                                                                              568 P-----NPIESLE-----KESAFKTKAEPISSSISNMELELNIHNSKAPKKNRL 611
                                                                                                                                                                                                                                                           292 KKGYCECCLQKYEDLETHLLSEOHRNFAQSNQYQVVDDIVSKLYFDFVEYEKDTPKKKRI 351
                                                                                                                                                                                                                                                                                                                 518 IKKADL----AVQK--TPEMINQGTNQTEQNGQVMNITNSGHENKTRGDSIQ---NEKN 567
                                                                                                                                                                                                                                                                                                                                                                 234 LQLTNMPFINYSIQKPCSPFDVDKPSSMQKQTQVKLRIQTDG--DKYGGTSIQLQLKEKK 291
                                                                                                                                                                                                                                                                                                                                                                                                                       461 FGKTYRKKASLPNLSHVTENLIIGAFVTEPQIIQERPLTNKLKR---KRRPTSGLHPEDF 517
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       183 KKELYLLKKSSTSVRDGGKRVGSGA------QKTRTGRLKKPFVKVEDMSQLYRPFY 233
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             423 SGSSEKI-DLLASDPHEALICKSERVH----SKSVESN-----IEDKI 460
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             124 DGSSFKSPDTVCLSRGKLLVEKAIKDHDFIPSNSILSNALSWGVKILHIDDIRYYIEQK- 182
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               379 IQK-VNEWFSRSDELLGSDDSHDGESESNÄKVADVLDVLNEV------DEY 422
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     323 GSKETCNDRRTPSTEKKVDLNADPLCERKE----WNKQKLPCSENPRDTEDVPWITLNSS 378
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT STREET: 4 Embarcadero Center, Suite 3400 CTIY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               64 LQKDIKDLGGRVEEFLSKDISYLISNKKEAKFAQTLGRISPVPSPESAYTAETTSPHPSH 123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16 GGIQVKNEKNRPSLK---SLKTDNRPEKSKCKPLWGKVFY------LDLPSVTISEK 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1.0%; Score 140.5; DB 1; Length 900; al Similarity 20.2%; Pred. No. 0.0024; al Similarity 206; Mismatches 239; Indels 205;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           94111-4187
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                                                                                                                                   ; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-825-886-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-825-886-20
Query Match 4.0%; Score 140.5; DB 2; Best Local Similarity 20.2%; Pred. No. 0.0024; Matches 139; Conservative 106; Mismatches 239;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sequence 20, Application US/08825886 Patent No. 5821328
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: KING, I
APPLICANT: FRIEDM
                                                                                                                                                                                                                                                                                                                    TELEFAX: (415) 494-877
TELEX: 910 277299
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 900 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: OSMAN, RICHARD A
REGISTRATION NUMBER: 36,627
REFERENCE/DOCKET NUMBER: A-
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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ZIP: 94111-4187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE:
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SZABO, Csilla
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FRIEDMAN, Lori
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(415) 494-8771
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TITLE OF INVENTION: CANCER
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
ADDRESSER: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT
STREET: 4 Embarcadero Center, Suite 3400
CITY: San Francisco
STATE: California
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Length 900;
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                                                                                                                                                                                                                                                                                                                                       Sequence 21, Application US/08425061 Patent No. 5622829 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                  APPLICANT: APPLICANT:
                                                                                                                             NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                  TITLE OF INVENTION: GENETIC MARKERS TITLE OF INVENTION: CANCER
                                                                                                                                                                                                                                                                    APPLICANT:
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COMPUTER READABLE FORM:
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850 SEL---DAQYLQNTFKVSKRQS-FAPFSN 874
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                   COUNTRY: USA
ZIP: 94111-4187
                                                      ADDRESSEE: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT STREET: 4 Embarcadero Center, Suite 3400 CITY: San Francisco STATE: California
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                                                    California
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SZABO, Csilla
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FRIEDMAN, Lori
OSTERMEYER, Beth
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; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-425-061-21
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INFORMATION FOR SEQ ID NO:
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NAME: OSMAN, RICHARD 36,627
REGISTRATION UNMBER: A-59
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 494-8770
TELEFAX: (415) 494-8771
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/425,061
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CLASSIFICATION: 435
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                                                                                                                                                                                                                         DDIRONFTQLPLHKNKQECILDISEHTLSENDLEELRVDHYKCNIQASVHVSDFSTDNSG 501
                                                                                                                                                                                                                                                                                                                   QEDDTTVKEQNFLYKETQETEKKLLFISEPIPHPSN---ELRGLNEKMS-NKCSMLSTAE 441
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GSKETCNDRRTPSTEKKVDLNADPLCERKE----WNKQKLPCSENPRDTEDVPWITLNSS
NKCVSQCAAFEN --- PKGLIHGCSKD -- NRNDTEGFKYPLGHEVN --- HSRETSIEMEE 849
                                                                                       EDPKDLMLSGERVLQTERSVESSSISLVPGTDYG-----TQESISLLEVSTLGKAKTEP
                                                                                                                                  SQPKQ---KSDTVLFPAKDLKEKDLHSIFTHDSGLITINSSQEHLTVQAKAPFHTPPEEP 558
                                                                                                                                                                                                                                                                                                                                                                  RRKSSTRHIHALELVVSRNLSPPNCTELQIDSCSSSEEIKKKKYNQMPVRHSRNLQLMEG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FGKTYRKKASLPNLSHVTENLIIGAFVTEPQIIQERPLTNKLKR----KRRPTSGLHPEDF
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                                          NEC-----DFKNMDSLPSGKIHRKVKIILGRNRKENL------EPNAEFDKRTEFITQEE 607
                                                                                                                                                                                 EFVNPS---LP----REE------KEEKLETVKV-----
                                                                                                                                                                                                                                                                           KEPATGAKKSN---KPNEQTSKR------HDSDTFPELKLTNAPGSFTKCSNTSELK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LQLTNMPFINYSIQKPCSPFDVDKPSSMQKQTQVKLRIQTDG--DKYGGTSIQLQLKEKK 291
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      914 amino acids
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Sequence 21, Application US/08825886
Patent No. 5821328
GENERAL INFORMATION:
APPLICANT: KING, Mary-Claire
APPLICANT: FRIEDMAN, Lori
APPLICANT: FRIEDMAN, Lori
APPLICANT: ROWELL, Sarah
APPLICANT: ROWELL, Sarah
APPLICANT: LYNGH, Eric
APPLICANT: LYNGH, Eric
APPLICANT: LYNGH, Eric
APPLICANT: LEE, Ming
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: OSMAN, Richard A
REGISTRATION NUMBER: 36,627
REFERENCE/DOCKET NUMBER: A-99563-3/DJB/RAO
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 494-8770
TELEPAX: (415) 494-8771
TELEPAX: (415) 494-8771
TELEPAX: 910 277299
INFORMATION FOR SEQ ID NO: 21:
SEQUENCE CHARACTERISTICS:
LENGTH: 914 amino acids
TYPE: amino acids
В
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MEDIUM TYPE: FORM:

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC COMPUTER:

SOFTWARE: PATENTION DATA:
APPLICATION UNMER: US/08/825,886

FILING DATE:
CLASSIFICATION NUMBER: US/08/825,886

FILING DATE:
APPLICATION NUMBER: 08/425,061

FILING DATE:
FILING DATE:
CLASSIFICATION NUMBER: 08/425,061

FILING DATE:
FILING DATE:
APPLICATION NUMBER: 08/425,061
                                                                                                                                                                                                                                                                                           Query Match 4.0%; Score 140.5; DB 2; Length 914; Best Local Similarity 20.2%; Pred. No. 0.0024; Matches 139; Conservative 106; Mismatches 239; Indels 205; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: GENETIC MARKERS FOR BREAST AND OVARIAN TITLE OF INVENTION: GANCER NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
ADDRESSEE: FLERR, HOHBACH, TEST, ALBRITTON & HERBERT STREET: 4 Embarcadero Center, Suite 3400
CITY: San Francisco
STATE: California
COUNTRY: USA
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                              124 DGSSFKSPDTVCLSRGKLLVEKAIKDHDFIPSNSILSNALSWGVKILHIDDIRYYIEQK- 182
                                                                                               379 IQK-VNEWFSRSDELLGSDDSHDGESESNAKVADVLDVLNEV------DEY 422
                                                                                                                                                                                              323 GSKETCNDRRTPSTEKKVDLNADPLCERKE----WNKQKLPCSENPRDTEDVPWITLNSS 378
423 SGSSEKI-DLLASDPHEALICKSERVH----SKSVESN------IEDKI 460
                                                                                                                              64 LQKDIKDLGGRVEEFLSKDISYLISNKKEAKFAQTLGRISPVPSPESAYTAETTSPHPSH 123 : | | | | | :: | :: | | :: |
                                                                                                                                                                                                                                               16 GGIQVKNEKNRPSLK---SLKTDNRPEKSKCKPLWGKVFY-----LDLPSVTISEK 63
                                                                                                                                                                                                                                                                                                          36;
         REFERENCE/DOCKET NUMBER: A-59563-3/DJB/RAO
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APPLICATION NUMBER: US/08/425,061 FILING DATE: CLASSIFICATION: 435 ATTORNEY_AGENT INFORMATION: ATTORNEY_AGENT INFORMATION: NAME: OSMAN, Richard A REGISTRATION NUMBER: 36,627 BEFERENCY_COCKET NUMBER: A-59563-3/DJB/RAO	, ,, ,, ,, ,, ,, ,, ,,
OMPUTER READABLE FO MEDIUM TYPE: Flop COMPUTER: IBM PC OPERATING SYSTEM: SOFTWARE: Patenti	
DENCE ADDRES EE: FLEHR, 4 Embarcac San Francis California : USA	
ABO, Cs ABO, Cs E, Ming TION: TION:	
Application: 622829 ORMATION: KING, PERIEDMA COSTERMI COSTERMI COSTERMI COSTERMI COSTERMI	
12	RESU
608 NRICSSPVQSLLDLPQTSEEKSEFLGFTS 636 :::- : : :	QQ dd
559 NECDFKNKDSLPSGKIHRKVKIILGRNRKENLEPNAEFDKRTEFITQEE 607 :	pb dy
502 SQPKQKSDTVLFPAKDLKEKDLHSIFTHDSGLITINSSQEHLTVQAKAPFHTPPEEP 558	Qy Db
442 DDIRQNFTQLPLHKNKQECILDISEHTLSENDLEELRVDHYKCNIQASVHVSDFSTDNSG 501	Qу
386 QEDDTTVKEONFLYKETQETEKKLLFISEPIPHPSNELRGLNEKMS-NKCSMLSTAE 441 :	Qу
352 KYSVGSLSPVSASVLKKTEQKEKVELQHISQKDC 385	Qy Db
292 KKGYCECCLOKYEDLETHLLSEOHRNFAQSNOYQVVDDIVSKLVFDFVEYEKDTPKKRI 351	Оу
234 LOLTNMPFINYSIOKPCSPEDVDKPSSMOKQTQVKLRIQTDGDKYGGTSIOLOLKEKK 291 ::::::::::::::::::::::::::::::::::::	Qy
183 KKELYLLKKSSTSVRDGGKRVGSGAQKTRTGRLKKPFVKVEDMSQLYRPFY 233	Фу

TELECOMMUNICATION INFORMATION:

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US-08-825-886-22
; Sequence 22, Application
; Patent No. 5821328
; GENERAL INFORMATION:
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Best Local Similarity
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SEQUENCE CHARACTERISTICS:
LENGTH: 1202 amino acids
    APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 1202 amino
TYPE: amino acid
STRANDEDNESS: sin
                                                                                                                                                                                               608 NRICSSPYQSLLDLFQTSEEKSEFLGFTS 636
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           672 KEPATGAKKSN----KPNEQTSKR-------HDSDTFPELKLTNAPGSFTKCSNTSELK 719
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           612 RRKSSTRHIHALELVVSRNLSPPNCTELQIDSCSSSEEIKKKKYNQMPVRHSRNLQLMEG 671
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     323 GSKETCNDRRTPSTEKKVDLNADPLCERKE----WNKQKLPCSENPRDTEDVPWITLNSS 378
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             124 DGSSFKSPDTVCLSRGKLLVEKAIKDHDFIPSNSILSNALSWGVKILHIDDIRYYIEQK- 182
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TELEX: 910 277299
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         P-----NPIESLE-----KESAFKTKAEPISSSISNMELELNIHNSKAPKKNRL
                                                                                                                                                                SEL---DAOYLONTFKVSKRQS-FAPFSN 874
                                                                                                                                                                                                                                                                                 NEC-----EPNAEFDKRTEFITQEE
                                                                                                                                                                                                                                                                                                                                                                 SQPKQ----KSDTVLFPAKDLKEKDLHSIFTHDSGLITINSSQEHLTVQAKAPFHTPPEEP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       IKKADL----AVQK--TPEMINQGTNQTEQNGQVMNITNSGHENKTKGDSIQ---NEKN 567
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LQLTIMPFINYSIQKPCSPFDVDKPSSMQKQTQVKLRIQTDG--DKYGGTSIQLQLKEKK 291
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SGSSEKI-DLLASDPHEALICKSERVH----SKSVESN------
                                                                                                                                                                                                                                             NKCVSQCAAFEN----PKGLIHGCSKD--NRNDTEGFKYPLGHEVN---HSRETSIEMEE
                                                                                                                                                                                                                                                                                                                                                                                                                                                 DDIRONFTQLPLHKNKQECILDISEHTLSENDLEELRVDHYKCNIQASVHVSDFSTDNSG 501
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 QEDDTTVKEQNFLYKETQETEKKLLFISEPIPHPSN---ELRGLNEKMS-NKCSMLSTAE 441
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     139;
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KING, Mary-Claire
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (415) 494-8700
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                                                          US/08825886
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 140.5; DB 1;
Pred. No. 0.0036;
06; Mismatches 239;
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Best Local Similarity
Matches 139; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (415) 494-877
TELEX: 910 277299
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
STRANDEDNESS: sin
TOPOLOGY: linear
MOLECULE TYPE: prot
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 94111-418/
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER: A-59563-3/DJB/RAO TELECOMMUNICATION INFORMATION: TELEPHONE: (415) 494-8700 TELEPAX: (415) 494-8771
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: OSMAN, Richard A
REGISTRATION NUMBER: 36,627
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/825,886
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: GENETIC TITLE OF INVENTION: CANCER UNMBER OF SEQUENCES: 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
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568
                                                                                                                                             461 FGKTYRKKASLPNLSHVTENLIIGAFVTEPQIIQERPLTNKLKR---KRRPTSGLHPEDF
                                                                                                                                                                                                                                         124 DGSSFKSPDTVCLSRGKLLVEKAIKDHDFIPSNSILSNALSWGVKILHIDDIRYYIEQK- 182
                                                                                                                                                                                                                                                                                                                                                                  323 GSKETCNDRRTPSTEKKVDLNADPLCERKE----WNKQKLPCSENPRDTEDVPWITLNSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET:
                                                                                                                                                                                                                                                                                                                            64 LQKDIKDLGGRVEEFLSKDISYLISNKKEAKFAQTLGRISPVPSPESAYTAETTSPHPSH 123
                                                                                                                                                                                                                                                                                                                                                                                                    16 GGIQVKNEKNRPSLK---SLKTDNRPEKSKCKPLWGKVFY------LDLPSVTISEK 63
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH:
                                                                                          LQLTNMPFINYSIQKPCSPFDVDKPSSMQKQTQVKLRIQTDG--DKYGGTSIQLQLKEKK 291
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                                 KKGYCECCLQKYEDLETHLLSEQHRNFAQSNQYQVVDDIVSKLVFDFVEYEKDTPKKKRI 351
                                                                                                                                                                                                                                                                                            IQK-VNEWFSRSDELLGSDDSHDGESESNAKVADVLDVLNEV---
                                                                       IKKADL----AVQK--TPEMINQGTNQTEQNGQVMNITNSGHENKTKGDSIQ---NEKN
                                                                                                                                                                             KKELYLLKKSSTSVRDGGKRVGSGA------QKTRTGRLKKPFVKVEDMSQLYRPFY 233
                                                                                                                                                                                                                   SGSSEKI-DLLASDPHEALICKSERVH----SKSVESN------IEDKI 460
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T: 4 Embarcadero Center, Suite
San Francisco
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1202 amino acids
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SZABO, Csilla
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FRIEDMAN, LC
OSTERMEYER,
                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ROWELL, Sarah
 -NPIESLE-----KESAFKTKAEPISSSISNMELELNIHNSKAPKKNRL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       single
                                                                                                                                                                                                                                                                                                                                                                                                                                                       4.0%; Score 140.5; DB
20.2%; Pred. No. 0.0036;
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                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                         239;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 1202;
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 611
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US-08-425-061-23; Sequence 23, Applicat.; Sequence 26, Applicat.; Patent No. 5622829; GENERAL INFORMATION:
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US-08-425-061-23
                                                                                                     CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: OSMAN, RICHARD A
REGISTRATION NUMBER: 36,627
REFERENCE/DOCKET NUMBER: A-59
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 494-8770
TELEPHONE: (415) 494-8771
TELEPHONE: (415) 494-8771
TELEPHONE: 1050 407-299
INFORMATION FOR SEQ ID NO: 23:
SEQUENCE CHARACTERISTICS:
LENGTH: 1363 amino acids
TYPE: Amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: KING, Mary-Claire APPLICANT: FRIEDMAN, LOTI APPLICANT: OSTERMEYER, Beth APPLICANT: ROWELL, Sarah APPLICANT: LYNCH, Eric APPLICANT: SZABO, CSilla APPLICANT: LEE, Ming
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: Patentin Rel
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT
STREET: 4 Embarcadero Center, Suite 3400
CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: LEE, Ming
TITLE OF INVENTION: GENETIC MARKERS FOR BREAST AND OVARIAN
TITLE OF INVENTION: CANCER
NUMBER OF SEQUENCES: 24
                   MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE:
STREET: 4
CITY: San
STATE: Cal
COUNTRY: U
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      672 KEPATGAKKSN----KPNEQTSKR-------HDSDTFPELKLTNAPGSFTKCSNTSELK 719
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  386 QEDDTTVKEQNFLYKETQETEKKLLFISEPIPHPSN---ELRGLNEKMS-NKCSMLSTAE 441
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                                            TYPE: amino acid
STRANDEDNESS: sig
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 94111-4187
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Best Local Similarity 20.2%; Pred. No. 0.0043;
Matches 139; Conservative 106; Mismatches 239;
850 SEL---DAQYLQNTFKVSKRQS-FAPFSN 874
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                                                            608 NRICSSPYQSLLDLFQTSEEKSEFLGFTS 636
                                                                                                                        799 NKCVSQCAAFEN----PKGLIHGCSKD--NRNDTEGFKYPLGHEVN---HSRETSIEMEE 849
                                                                                                                                                                                 559 NEC-----DFKNMDSLPSGKIHRKVKIILGRNRKENL-----EPNAEFDKRTEFITQEE 607
                                                                                                                                                                                                                                             745 EDPKDLMLSGERVLQTERSVESSSISLVPGTDYG-----TQESISLLEVSTLGKAKTEP 798
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              568 P-----NPIESLE-----KESAFKTKAEPISSSISNMELELNIHNSKAPKKNRL 611
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   292 KKGYCECCLQKYEDLETHLLSEQHRNFAQSNQYQVVDDIVSKLVFDFVEYEKDTPKKKRI 351
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     234 LQLTNMPFINYSIQKPCSPFDVDKPSSMQKQTQVKLRIQTDG--DKYGGTSIQLQLKEKK 291
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      124 DGSSFKSPDTVCLSRGKLLVEKAIKDHDFIPSNSILSNALSWGVKILHIDDIRYYIEQK- 182
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   379 IQK-VNEWFSRSDELLGSDDSHDGESESNAKVADVLDVLNEV-------DEY 422
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16 GGIQVKNEKNRPSLK---SLKTDNRPEKSKCKPLWGKVFY-----LDLPSVTISEK 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           64 LQKDIKDLGGRVEEFLSKDISYLISNKKEAKFAQTLGRISPVPSPESAYTAETTSPHPSH 123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       IKKADL----AVQK--TPEMINQGTNQTEQNGQVMNITNSGHENKTKGDSIQ---NEKN 567
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4.0%; Score 140.5; DB 1; Length 1363; 20.2%; Pred. No. 0.0043;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels 205; Gaps
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RESULT 15
US-08-825-886-23
US-08-825-886-23
Sequence 23, Application US/08825886
Patent No. 5821328
GEMERAL INFORMATION:
APPLICANT: KING, MATY-Claire
APPLICANT: FRIEDMAN, Lori
APPLICANT: CSTERREYER, Beth
APPLICANT: STACH, Eric
APPLICANT: STACH, Eric
APPLICANT: STACH, Cailla
APPLICANT: LEE, Ming
TITLE OF INVENTION: GENETIC MARKERS FOR BREAST AND OVARIAN
TITLE OF INVENTION: CANCER
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
ADDRESSEET: 4 Embarcadero Center, Smite 3400

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; LENGTH: 1363 amino aci
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-825-886-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 4.0%; Score 140.5; DB 2; Length 1363; Best Local Similarity 20.2%; Pred. No. 0.0043; Matches 139; Conservative 106; Mismatches 239; Indels 205; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENCTH: 1363 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: OSMAN, Richard A
REGISTRATION NUMBER: 36,627
REFERENCE/DOCKET NUMBER: A-59563-3/DJB/RAO
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 494-8771
TELEPHONE: (415) 494-8771
TELEX: 910 277299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      323 GSKETCNDRRTPSTEKKVDLNADPLCERKE----WNKQKLPCSENPRDTEDVPWITLNSS 378
                                                                                                                                                                                                                                                                                                                                                                                                       292 KKGYCECCLQKYEDLETHLLSEQHRNFAQSNQYQVVDDIVSKLVFDFVEYEKDTPKKKRI 351
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              518 IKKADL-----AVQK--TPEMINQGTNQTEQNGQVMNITNSGHENKTKGDSIQ---NEKN 567
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         461 FGKTYRKKASLPNLSHVTENLIIGAFVTEPQIIQERPLTNKLKR---KRRPTSGLHPEDF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        379 IQK-VNEWFSRSDELLGSDDSHDGESESNAKVADVLDVLNEV------DEY 422
                                                          442 DDIRONFTOLPLHKNKOECILDISEHTLSENDLEELRVDHYKCNIQASVHVSDFSTDNSG 501
                                                                                                                       672 KEPATGAKKSN---KPNEQTSKR-----HDSDTFPELKLTNAPGSFTKCSNTSELK 719
                                                                                                                                                                                                                                               612 RRKSSTRHIHALELVVSRNLSPPNCTELQIDSCSSSEEIKKKKYNQMPVRHSRNLQLMEG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                183 KKELYLLKKSSTSVRDGGKRVGSGA------QKTRTGRLKKPFVKVEDMSQLYRPFY 233
720 EFVNPS---LP----REE------KEEKLETVKV------SNNA 744
                                                                                                                                                                                 386 QEDDTTVKEQNFLYKETQETEKKLLFISEPIPHPSN----ELRGLNEKMS-NKCSMLSTAE 441
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  64 LQKDIKDLGGRVEEFLSKDISYLISNKKEAKFAQTLGRISPVPSPESAYTAETTSPHPSH 123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16 GGIQVKNEKNRPSLK---SLKTDNRPEKSKCKPLWGKVFY------LDLPSVTISEK 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                          P-----NPIESLE-----KESAFKTKAEPISSSISNMELELNIHNSKAPKKNRL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LQLTNMPFINYSIQKPCSPFDVDKPSSMQKQTQVKLRIQTDG--DKYGGTSIQLQLKEKK 291
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SGSSEKI-DLIASDPHEALICKSERVH----SKSVESN-----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  94111-4187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SS: single
linear
                                                                                                                                                                                                                                                                                                         -----KKTEQKEKVELQHISQKDC 385
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           08/425,061
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                                            608 NRICSSPYOSILDLFOTSEEKSEFLGFTS 636
                                                                                                                                       559 NEC-----DFKNMDSLPSGKIHRKVKIILGRNRKENL-----EPNAEFDKRTEFITQEE 607
                                                                                                                                                                                  745 EDPKDLMLSGERVLQTERSVESSSISLVPGTDYG-----TQESISLLEVSTLGKAKTEP
                                                                                                                                                                                                                            502 SQPKQ----KSDTVLFPAKDLKEKDLHSIFTHDSGLITINSSQEHLTVQAKAPFHTPPEEP
SEL---DAOYLONTFKVSKROS-FAPFSN 874
                                                                                           NKCVSQCAAFEN----PKGLIHGCSKD--NRNDTEGFKYPLGHEVN---HSRETSIEMEE
                                                                                                                                                                                    798
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